

=====

Sequence Listing could not be accepted due to errors.
See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2010; month=11; day=3; hr=10; min=53; sec=1; ms=998;]

=====

Reviewer Comments:

<210> 1

<211> 7

<212> PRT

<213> Gymnea sylvestre

<400> 1

Asn Gly Ser Phe Ser Gly Phe

1 5

The above sequence id# 1 is invalid, please delete spacing between the amino acids and the numbering. This error is seen globally throughout the sequences.

Application No: 10579655 Version No: 4.0

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (5)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (7)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (8)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (8)
W 402	Undefined organism found in <213> in SEQ ID (9)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (9)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (9)
W 402	Undefined organism found in <213> in SEQ ID (10)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (10)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (10)

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (11)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (11)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (11)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (11)
W 402	Undefined organism found in <213> in SEQ ID (12)
E 355	Empty lines found between the amino acid numbering and the proteins
E 321	No. of Bases conflict, this line has no nucleotides SEQID (12) POS (0)
W 402	Undefined organism found in <213> in SEQ ID (13)
W 402	Undefined organism found in <213> in SEQ ID (14)
W 402	Undefined organism found in <213> in SEQ ID (15)
W 402	Undefined organism found in <213> in SEQ ID (16)
W 402	Undefined organism found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22)
W 402	Undefined organism found in <213> in SEQ ID (23)

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (24)
W 402	Undefined organism found in <213> in SEQ ID (25)
W 402	Undefined organism found in <213> in SEQ ID (26)
W 402	Undefined organism found in <213> in SEQ ID (28)
W 402	Undefined organism found in <213> in SEQ ID (29)
W 402	Undefined organism found in <213> in SEQ ID (30) This error has occurred more than 20 times, will not be displayed
W 213	Artificial or Unknown found in <213> in SEQ ID (77)
W 213	Artificial or Unknown found in <213> in SEQ ID (78)
W 213	Artificial or Unknown found in <213> in SEQ ID (79)
W 213	Artificial or Unknown found in <213> in SEQ ID (80)
W 213	Artificial or Unknown found in <213> in SEQ ID (253)
W 213	Artificial or Unknown found in <213> in SEQ ID (254)
W 213	Artificial or Unknown found in <213> in SEQ ID (255)
W 213	Artificial or Unknown found in <213> in SEQ ID (256)
W 213	Artificial or Unknown found in <213> in SEQ ID (257)
W 213	Artificial or Unknown found in <213> in SEQ ID (258)
W 213	Artificial or Unknown found in <213> in SEQ ID (259) This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

<110> Sanofi Pasteur, Inc.

<120> METHODS FOR PURIFYING PERTUSSIS TOXIN AND PEPTIDES USEFUL
THEREFOR

<130> API-03-15

<140> 10579655

<141> 2010-10-25

<150> 60/523,881

<151> 2003-11-20

<150> PCT/US2004/038700

<151> 2004-11-18

<160> 415

<170> PatentIn version 3.5

<210> 1

<211> 7

<212> PRT

<213> *Gymnea sylvestre*

<400> 1

Asn Gly Ser Phe Ser Gly Phe

1 5

<210> 2

<211> 7

<212> PRT

<213> *Gymnea sylvestre*

<400> 2

Asn Gly Ser Phe Ser Gly Cys

1 5

<210> 3

<211> 7

<212> PRT

<213> *Gymnea sylvestre*

<400> 3

Asp Gly Ser Phe Ser Gly Phe

1 5

<210> 4

<211> 7

<212> PRT

<213> Gymnea sylvestre

<220>

<221> MISC_FEATURE

<222> (1) .. (7)

<223> Xaa is any amino acid

<400> 4

Xaa Gly Ser Phe Ser Gly Xaa

1 5

<210> 5

<211> 30

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 5

Arg Ser Ser His Cys Arg His Arg Asn Cys His Thr Ile Thr Arg Gly

1 5 10 15

Asn Met Arg Ile Glu Thr Pro Asn Asn Ile Arg Lys Asp Ala

20

25

30

<210> 6

<211> 29

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 6

Ser Thr Met Asn Thr Asn Arg Met Asp Ile Gln Arg Leu Met Thr Asn

1

5

10

15

His Val Lys Arg Asp Ser Ser Pro Gly Ser Ile Asp Ala

20

25

<210> 7

<211> 30

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 7

Arg Ser Asn Val Ile Pro Leu Asn Glu Val Trp Tyr Asp Thr Gly Trp

1 5 10 15

Asp Arg Pro His Arg Ser Arg Leu Ser Ile Asp Asp Asp Ala

20 25 30

<210> 8

<211> 30

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 8

Arg Ser Trp Arg Asp Thr Arg Lys Leu His Met Arg His Tyr Phe Pro

1 5 10 15

Leu Ala Ile Asp Ser Tyr Trp Asp His Thr Leu Arg Asp Ala

20 25 30

<210> 9

<211> 34

<212> PRT

<213> *Gymnea sylvestre*

<400> 9

Ser Gly Cys Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val

1

5

10

15

Cys Cys Glu Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr

20

25

30

Cys Gly

<210> 10

<211> 34

<212> PRT

<213> *Gymnea sylvestre*

<400> 10

Ser Gly Cys Val Lys Lys Asp Glu Leu Cys Glu Leu Ala Val Asp Glu

1

5

10

15

Cys Cys Glu Pro Leu Glu Cys Phe Gln Met Gly His Gly Phe Lys Arg

20

25

30

Cys Gly

<210> 11

<211> 35

<212> PRT

<213> *Gymnea sylvestre*

<400> 11

Ser Gly Cys Val Lys Lys Asp Glu Leu Cys Ser Gln Ser Val Pro Met

1

5

10

15

Cys Cys Glu Pro Leu Glu Cys Lys Trp Phe Asn Glu Asn Tyr Gly Ile

20

25

30

Cys Gly Ser

35

<210> 12

<211> 34

<212> PRT

<213> *Gymnea sylvestre*

<400> 12

Ser Gly Cys Val Lys Lys Asp Glu Leu Cys Glu Leu Ala Ile Asp Glu

1 5 10 15

Cys Cys Glu Pro Leu Glu Cys Thr Lys Gly Asp Leu Gly Phe Arg Lys

20 25 30

Cys Gly

<210> 13

<211> 35

<212> PRT

<213> *Gymnea sylvestre*

<400> 13

Gln Gln Cys Val Lys Lys Asp Glu Leu Cys Ile Pro Tyr Tyr Leu Asp

1 5 10 15

Cys Cys Glu Pro Leu Glu Cys Lys Lys Val Asn Trp Trp Asp His Lys

20

25

30

Cys Ile Gly

35

<210> 14

<211> 31

<212> PRT

<213> Gymnea sylvestre

<220>

<221> MISC_FEATURE

<222> (9)..(30)

<223> Xaa is any amino acid

<400> 14

Cys Val Lys Lys Asp Glu Leu Cys Xaa Xaa Xaa Xaa Xaa Cys Cys

1

5

10

15

Glu Pro Leu Glu Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys

20

25

30

<210> 15

<211> 141

<212> DNA

<213> *Gymnea sylvestre*

<220>

<221> misc_feature

<222> (49)..(113)

<223> n is a, g, t or c

<400> 15

agtggctcaa gctcaggatc aggctgcgtc aagaaagacg agctctgcn snnsnnnsns 60

nnsnnnstgct gtgagccccct cgagtgcnn nnsnnsnnsn nsnnsnnnsnn snnstgcggc 120

agcggcagtt ctgggtctag c 141

<210> 16

<211> 84

<212> DNA

<213> *Gymnea sylvestre*

<400> 16

taatacggact cactataggg acaattacta tttacaatta caatgcacca tcaccatcac 60

catagtggct caagctcagg atca

84

<210> 17

<211> 44

<212> DNA

<213> *Gymnea sylvestre*

<400> 17

ttttaatacg cggatgctac taggctagac ccagaactgc cgct

44

<210> 18

<211> 10

<212> RNA

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 18

uagcggaugc

10

<210> 19

<211> 53

<212> PRT

<213> Artificial sequence

<220>

<223> Synthetic sequence, no source organism

<220>

<221> MISC_FEATURE

<222> (18) .. (43)

<223> Xaa is any amino acid

<400> 19

Thr Met Val Met Gly Arg Gly Ser His His His His His Ala Arg

1 5 10 15

Ser Xaa Xaa

20 25 30

Xaa Asp Ala Asn Ala Pro

35 40 45

Lys Ala Ser Ala Ile

50

<210> 20

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 20

His His His His His His

1 5

<210> 21

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 21

Asp Ala Asn Ala Pro Lys

1 5

<210> 22

<211> 127

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic sequence, no source organism

<220>

<221> MISC_FEATURE

<222> (28) .. (105)

<223> S represents C or G, and N represents A, T, G or C

<400> 22

agcggatgcc ttccggagcgt tagcgtcsnn snnsnnsnns nnnsnnsnnsn nnnsnnnnnn 60

snnsnnsnns nnnsnnnnn nsnnnnnnn snnsnnsnns nnnsnnagatc tagcatgatg 120

atgatga 127

<210> 23

<211> 81

<212> DNA

<213> Gymnea sylvestre

<400> 23

taatacgact catagggaca attactattt acaattacaa tgggacgtgg ctcacatcat 60

catcatcatc atgctagatc t 81

<210> 24

<211> 32

<212> DNA

<213> *Gymnea sylvestre*

<400> 24

aat taaaat a g cggatgcctt cggagcgtta g c 32

<210> 25

<211> 18

<212> DNA

<213> Bacteriophage M13

<400> 25

t g taaaac g a cggccagt 18

<210> 26

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 26

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

Val Lys Lys Asp Glu Leu Cys Ala Gly Ser Val Gly His Cys Cys Glu

20

25

30

Pro Leu Glu Cys Leu Arg Arg Phe Leu Asn Leu Arg Trp Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 27

<211> 54

<212> PRT

<213> *Gymnema sylvestre*

<400> 27

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1

5

10

15

Val Lys Lys Asp Glu Leu Cys Ile Val Met Arg Ala Pro Cys Cys Glu

20

25

30

Pro Leu Glu Cys Leu Arg Arg Tyr Met Leu Lys His Met Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 28

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 28

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1

5

10

15

Val Lys Lys Asp Glu Leu Cys Lys Ala Phe Arg Tyr Ser Cys Cys Glu

20

25

30

Pro Leu Glu Cys Leu Arg Lys Trp Leu Lys Ala Arg Phe Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 29

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 29

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Leu Arg Ser Ser Ile Asp Cys Cys Glu

20 25 30

Pro Leu Glu Cys Leu Tyr Lys Trp Met Gln Arg Arg Leu Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 30

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 30

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Trp Pro Arg Arg His Lys Cys Cys Glu

20 25 30

Pro Leu Glu Cys Leu Leu Glu Met Leu Glu Arg Lys Arg Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 31

<211> 53

<212> PRT

<213> Gymnea sylvestre

<400> 31

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Met Ser Met Ala Cys Val Cys Cys Glu

20

25

30

Pro Leu Glu Cys Lys Tyr His Gly Tyr Phe Trp Leu Cys Gly Ser Gly

35

40

45

Ser Ser Gly Ser Ser

50

<210> 32

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 32

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1

5

10

15

Val Lys Lys Asp Glu Leu Cys Ala Val Trp Phe Asp Val Cys Cys Glu

20

25

30

Pro Leu Glu Cys Thr Tyr Gln Ser Gly Tyr Tyr Trp Leu Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 33

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 33

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1

5

10

15

Val Lys Lys Asp Glu Leu Cys Glu Pro Trp Tyr Trp Arg Cys Cys Glu

20

25

30

Pro Leu Glu Cys Val Tyr Thr Ser Gly Tyr Tyr Ser Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

<210> 34

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 34

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 35

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 35

Met His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Val Phe Tyr Phe Pro Asn Cys Cys Glu

20 25 30

Pro Leu Glu Cys Arg Trp Val Asn Asp Asn Tyr Gly Trp Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 36

<211> 53

<212> PRT

<213> *Gymnea sylvestre*

<400> 36

Met His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Met Ser Met Ala Cys Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Lys Tyr His Gly Tyr Phe Trp Leu Cys Gly Ser Gly

35 40 45

Ser Ser Gly Ser Ser

50

<210> 37

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 37

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Thr Thr Ala Ser Lys Ser Cys Cys Glu

20 25 30

Pro Leu Glu Cys Lys Trp Thr Asn Glu His Phe Gly Thr Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 38

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 38

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1

5

10

15

Val Lys Lys Asp Glu Leu Cys Ser Gln Ser Val Pro Met Cys Cys Glu

20

25

30

Pro Leu Glu Cys Lys Trp Phe Asn Glu Asn Tyr Gly Ile Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 39

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 39

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 40

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 40

Met His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Leu Gly His Gly Leu Gly Tyr Ala Tyr Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 41

<211> 53

<212> PRT

<213> *Gymnea sylvestre*

<400> 41

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Met Trp Ser Arg Glu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Tyr Tyr Thr Gly Trp Tyr Trp Ala Cys Gly Ser Gly

35 40 45

Ser Ser Gly Ser Ser

50

<210> 42

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 42

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Glu Leu Ala Val Asp Glu Cys Cys Glu

20

25

30

Pro Leu Glu Cys Phe Gln Met Gly His Gly Phe Lys Arg Cys Gly Ser

35

40

45

Gly Ser Ser Gly Ser Ser

50

<210> 43

<211> 54